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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/511,939

DATE: 04/24/2002
TIME: 14:20:55

p6

Input Set : A:\seqlist.ST25.txt
Output Set: N:\CRF3\04242002\I511939.raw

```
    3 <110> APPLICANT: Tomlinson, Ian M
    4        Winter, Gregory
    6 <120> TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different
Ligands
    8 <130> FILE REFERENCE: 8039/1070
   10 <140> CURRENT APPLICATION NUMBER: US 09/511,939
C-->11 <141> CURRENT FILING DATE: 2002-04-10
   13 <150> PRIOR APPLICATION NUMBER: GB 9722131.1
   14 <151> PRIOR FILING DATE: 1997-10-20
   16 <150> PRIOR APPLICATION NUMBER: US 60/065,248
   17 <151> PRIOR FILING DATE: 1997-11-13
   19 <150> PRIOR APPLICATION NUMBER: US 60/066,729
   20 <151> PRIOR FILING DATE: 1997-11-21
   22 <150> PRIOR APPLICATION NUMBER: PCT/GB98/03135
   23 <151> PRIOR FILING DATE: 1998-10-20
   25 <160> NUMBER OF SEQ ID NOS: 350
   27 <170> SOFTWARE: PatentIn version 3.1
   29 <210> SEQ ID NO: 1
   30 <211> LENGTH: 720
   31 <212> TYPE: DNA
   32 <213> ORGANISM: Homo sapiens
   34 <400> SEQUENCE: 1
   35 gaggtgcagc tgttggagtc tggggggaggc ttggtacagc ctggggggtc cctgagactc      60
   37 tcctgtgcag cctctggatt cacctttagc agctatgcca tgagctgggt ccgccaggct     120
   39 ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac     180
   41 gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat     240
   43 ctgcaaatga acagcctgag agccgaggac acggccgtat attactgtgc gaaaagttat     300
   45 ggtgcttttg actactgggg ccagggaacc ctggtcaccg tctcgagcgg tggaggcggt     360
   47 tcaggcggag gtggcagcgg cggtggcggg tcgacggaca tccagatgac ccagtctcca     420
   49 tcctccctgt ctgcatctgt aggagacaga gtcaccatca cttgccgggc aagtcagagc     480
   51 attagcagct atttaaattg gtatcagcag aaaccaggga aagcccctaa gctcctgatc     540
   53 tatgctgcat ccagtttgca aagtggggtc ccatcaaggt tcagtggcag tggatctggg     600
   55 acagatttca ctctcaccat cagcagtctg caacctgaag attttgcaac ttactactgt     660
   57 caacagagtt acagtacccc taatacgttc ggccaaggga ccaaggtgga aatcaaacgg     720
   60 <210> SEQ ID NO: 2
   61 <211> LENGTH: 240
   62 <212> TYPE: PRT
   63 <213> ORGANISM: Homo sapiens
   65 <400> SEQUENCE: 2
   67 Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
   68 1               5                   10                  15
   71 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
   72             20                  25                  30
   75 Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
```

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Input Set : A:\seqlist.ST25.txt
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```
76            35                        40                        45
79  Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
80        50                     55                  60
83  Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
84  65                     70                  75                  80
87  Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
88                     85                  90                  95
91  Ala Lys Ser Tyr Gly Ala Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val
92             100                 105                 110
95  Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly
96             115                 120                 125
99  Gly Gly Ser Thr Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser
100     130                 135                 140
103 Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser
104 145                 150                 155                 160
107 Ile Ser Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro
108                 165                 170                 175
111 Lys Leu Leu Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser
112             180                 185                 190
115 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser
116             195                 200                 205
119 Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr
120     210                 215                 220
123 Ser Thr Pro Asn Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
124 225                 230                 235                 240
127 <210> SEQ ID NO: 3
128 <211> LENGTH: 7
129 <212> TYPE: DNA
130 <213> ORGANISM: Artificial Sequence
132 <220> FEATURE:
133 <223> OTHER INFORMATION: Artificial DVT variable codons used to introduce sequence
diversi
134       ty.
136 <220> FEATURE:
137 <221> NAME/KEY: misc_feature
138 <222> LOCATION: (1)..(7)
139 <223> OTHER INFORMATION: Artificial DVT variable codons used to introduce sequence
diversi
140       ty
143 <400> SEQUENCE: 3
144 agtagct
147 <210> SEQ ID NO: 4
148 <211> LENGTH: 7
149 <212> TYPE: DNA
150 <213> ORGANISM: Artificial Sequence
152 <220> FEATURE:
153 <223> OTHER INFORMATION: Artificial DVC variable codon used to introduce sequence
diversit
154       y.
156 <220> FEATURE:
157 <221> NAME/KEY: misc_feature
158 <222> LOCATION: (1)..(7)
```

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```
     159 <223> OTHER INFORMATION: Artificial DVC variable codon used to introduce sequence
diversit
     160      y.
     163 <400> SEQUENCE: 4                                                              7
     164 agtagcc
     167 <210> SEQ ID NO: 5
     168 <211> LENGTH: 7
     169 <212> TYPE: DNA
     170 <213> ORGANISM: Artificial Sequence
     172 <220> FEATURE:
     173 <223> OTHER INFORMATION: Artificial DVY codon used to introduce sequence variation.
     175 <220> FEATURE:
     176 <221> NAME/KEY: misc_feature
     177 <222> LOCATION: (1)..(7)
     178 <223> OTHER INFORMATION: Artificial DVY codon used to introduce sequence variation.
     181 <400> SEQUENCE: 5                                                              7
     182 agtagcy
     185 <210> SEQ ID NO: 6
     186 <211> LENGTH: 15
     187 <212> TYPE: PRT
     188 <213> ORGANISM: Artificial Sequence
     190 <220> FEATURE:
     191 <223> OTHER INFORMATION: Example of artificial linker sequence useful between VH and
VL do
     192      mains of scFV.
     194 <220> FEATURE:
     195 <221> NAME/KEY: MISC_FEATURE
     196 <223> OTHER INFORMATION: Example of artificial linker sequence useful between VL and
VH do
     197      mains of scFv.
     200 <400> SEQUENCE: 6
     202 Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
     203 1               5                   10                  15
     206 <210> SEQ ID NO: 7
     207 <211> LENGTH: 5
     208 <212> TYPE: PRT
     209 <213> ORGANISM: Homo sapiens
     211 <400> SEQUENCE: 7
     213 Ser Tyr Ala Met Ser
     214 1               5
     217 <210> SEQ ID NO: 8
     218 <211> LENGTH: 17
     219 <212> TYPE: PRT
     220 <213> ORGANISM: Homo sapiens
     222 <400> SEQUENCE: 8
     224 Ile Ile Gly Ser Glu Gly Trp Pro Thr Ile Tyr Ala Asp Ser Val Lys
     225 1               5                   10                  15
     228 Gly
     232 <210> SEQ ID NO: 9
     233 <211> LENGTH: 7
     234 <212> TYPE: PRT
     235 <213> ORGANISM: Homo sapiens
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/511,939

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Input Set : A:\seqlist.ST25.txt
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```
237 <400> SEQUENCE: 9
239 Gly Gly Ser Met Phe Asp Tyr
240 1               5
243 <210> SEQ ID NO: 10
244 <211> LENGTH: 11
245 <212> TYPE: PRT
246 <213> ORGANISM: Homo sapiens
248 <400> SEQUENCE: 10
250 Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn
251 1               5                   10
254 <210> SEQ ID NO: 11
255 <211> LENGTH: 7
256 <212> TYPE: PRT
257 <213> ORGANISM: Homo sapiens
259 <400> SEQUENCE: 11
261 Arg Ala Ser Ser Leu Gln Ser
262 1               5
265 <210> SEQ ID NO: 12
266 <211> LENGTH: 9
267 <212> TYPE: PRT
268 <213> ORGANISM: Homo sapiens
270 <400> SEQUENCE: 12
272 Gln Gln Ser Ser Asn Thr Pro Tyr Thr
273 1               5
276 <210> SEQ ID NO: 13
277 <211> LENGTH: 5
278 <212> TYPE: PRT
279 <213> ORGANISM: Homo sapiens
281 <400> SEQUENCE: 13
283 Ala Tyr Ala Met Thr
284 1               5
287 <210> SEQ ID NO: 14
288 <211> LENGTH: 17
289 <212> TYPE: PRT
290 <213> ORGANISM: Homo sapiens
292 <400> SEQUENCE: 14
294 Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys
295 1               5                   10                  15
298 Gly
302 <210> SEQ ID NO: 15
303 <211> LENGTH: 7
304 <212> TYPE: PRT
305 <213> ORGANISM: Homo sapiens
307 <400> SEQUENCE: 15
309 Lys Ala Ser Ser Phe Asp Tyr
310 1               5
313 <210> SEQ ID NO: 16
314 <211> LENGTH: 11
315 <212> TYPE: PRT
```

```
316 <213> ORGANISM: Homo sapiens
318 <400> SEQUENCE: 16
320 Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn
321 1               5                   10
324 <210> SEQ ID NO: 17
325 <211> LENGTH: 7
326 <212> TYPE: PRT
327 <213> ORGANISM: Homo sapiens
329 <400> SEQUENCE: 17
331 Ala Ala Ser Ser Leu Gln Ser
332 1               5
335 <210> SEQ ID NO: 18
336 <211> LENGTH: 9
337 <212> TYPE: PRT
338 <213> ORGANISM: Homo sapiens
340 <400> SEQUENCE: 18
342 Gln Gln Ser Tyr Ser Thr Pro Ser Thr
343 1               5
346 <210> SEQ ID NO: 19
347 <211> LENGTH: 5
348 <212> TYPE: PRT
349 <213> ORGANISM: Homo sapiens
351 <400> SEQUENCE: 19
353 Ser Tyr Ala Met Ser
354 1               5
357 <210> SEQ ID NO: 20
358 <211> LENGTH: 17
359 <212> TYPE: PRT
360 <213> ORGANISM: Homo sapiens
362 <400> SEQUENCE: 20
364 Leu Ile Ser Pro Leu Gly Lys Asp Thr Ser Tyr Ala Asp Ser Val Lys
365 1               5                   10                  15
368 Gly
372 <210> SEQ ID NO: 21
373 <211> LENGTH: 7
374 <212> TYPE: PRT
375 <213> ORGANISM: Homo sapiens
377 <400> SEQUENCE: 21
379 Arg Ala Gly Ile Phe Asp Tyr
380 1               5
383 <210> SEQ ID NO: 22
384 <211> LENGTH: 11
385 <212> TYPE: PRT
386 <213> ORGANISM: Homo sapiens
388 <400> SEQUENCE: 22
390 Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn
391 1               5                   10
394 <210> SEQ ID NO: 23
395 <211> LENGTH: 7
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 04/24/2002
PATENT APPLICATION: US/09/511,939   TIME: 14:20:56

    Input Set : **A:\seqlist.ST25.txt**
    Output Set: **N:\CRF3\04242002\I511939.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:320; N Pos. 23,24,29,30,32,33,38,39,41,42,44,45,50,51
Seq#:323; N Pos. 23,24,26,27,29,30,32,33
Seq#:326; N Pos. 21,22,30,31
Seq#:329; N Pos. 23,24,29,30,32,33,35,36,38,39
Seq#:332; N Pos. 21,22,27,28,33,34
Seq#:335; N Pos. 23,24,26,27,29,30,32,33
Seq#:338; N Pos. 21,22,27,28,30,31,33,34
Seq#:341; N Pos. 23,24